

Please amend the application as follows:

In the Claims:

Please cancel claims 141-205 and 219-247 without prejudice.

Please replace claims 46 and 61 with the following amended claims.

46. (New) An isolated polypeptide comprising an amino acid sequence 90% or more identical to an amino acid sequence selected from the group consisting of:

- C¹
- (a) amino acid residues 1 to 300 of SEQ ID NO:2;
 - (b) amino acid residues 2 to 300 of SEQ ID NO:2;
 - (c) amino acid residues 31 to 300 of SEQ ID NO:2; and
 - (d) amino acid residues 31 to 283 of SEQ ID NO:2;

wherein said polypeptide binds Fas ligand.

61. (New) An isolated polypeptide comprising an amino acid sequence 90% or more identical to an amino acid sequence selected from the group consisting of:

- C²
- (a) the amino acid sequence of the full-length polypeptide encoded by the cDNA contained in ATCC Deposit No. 97810.
 - (b) the amino acid sequence of the full-length polypeptide excluding the amino-terminal methionine encoded by the cDNA contained in ATCC Deposit No. 97810.
 - (c) the amino acid sequence of the mature polypeptide encoded by the cDNA contained in ATCC Deposit No. 97810; and
 - (d) the amino acid sequence of the extracellular domain of the polypeptide encoded by the cDNA clone contained in ATCC Deposit No. 97810;

wherein said polypeptide binds Fas ligand.